

**AMENDMENT**

**In the Specification**

Kindly replace the first full paragraph on page 13 of the specification with the following amended paragraph.

-- In order to determine the entire sequence, the enzyme was digested with trypsin and GluC. GluC is an endoproteinase (*Staphylococcus aureus* Protease V8) which is a serine proteinase, and which selectively cleaves peptide bonds C-terminal to glutamic acid residues (see Drapeau, G.R., Boily, Y. and Houmard, J. (1972); Purification and properties of an extracellular protease of *Staphylococcus aureus*. *J. Biol. Chem.*, 247, 6720-6726). The separation of the resulting peptides was effected by HPLC (reversed phase). The N-terminal sequence and the peptide fractions from the digestion of the BTA-hydrolase were analyzed by Edman degradation in an Applied Biosystems 473A Sequencer (gas-phase mode) or in a 494A Procise HT Sequencer (gas-phase and pulsed-liquid mode) using standard programs from the manufacturer.--